STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
   # ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information
Center (STIC) detected errors when processing the following computer readable
form:

Application Serial Number: _10/544,284B_

Source: _IFWP_

Date Processed by STIC: _2/10/07_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE
   APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
   NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
   http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is
a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the
United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission
   User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
   Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: _10/544,284B_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____Wrapped Nucleics
Wrapped Aminos

The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2____Invalid Line Length

The rules require that a line **not exceed 72** characters in length. This includes white spaces.

3____Misaligned Amino
Numbering

The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.

4____Non-ASCII

The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5____Variable Length

Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____PatentIn 2.0
"bug"

A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7____Skipped Sequences
(OLD RULES)

Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped
Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8____Skipped Sequences
(NEW RULES)

Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence.
<210> sequence id number
<400> sequence id number
000

9____Use of n's or Xaa's
(NEW RULES)

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of **n or Xaa**, and which residue n or Xaa represents.

10____Invalid <213>
Response

Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence. (see item 11 below)

11____Use of <220>

Sequence(s)____missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules

12____PatentIn 2.0
"bug"

Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____Misuse of n/Xaa

"n" can **only** represent a single <u>nucleotide</u>; "Xaa" can **only** represent a single <u>amino acid</u>

IFWP

RAW SEQUENCE LISTING   DATE: 02/10/2007
PATENT APPLICATION: US/10/544,284B   TIME: 09:53:52

Input Set : A:\corrected 70235USPCT.ST25.txt
Output Set: N:\CRF4\02102007\J544284B.raw

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 3 <110> APPLICANT: Brown, Devon
 4        Campos, Manuel
 5        Dalmia, Bipin
 6        Demarest, Stephen
 7        Hansen, Genevieve
 8        Heifetz, Peter B.
10 <120> TITLE OF INVENTION: Expression in plants of antibodies against enterotoxigenic
11        Escherichia coli
13 <130> FILE REFERENCE: 70235USPCT
15 <140> CURRENT APPLICATION NUMBER: 10/544,284B
16 <141> CURRENT FILING DATE: 2005-08-02
18 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/001427
19 <151> PRIOR FILING DATE: 2004-02-16
21 <150> PRIOR APPLICATION NUMBER: US 60/448,429
22 <151> PRIOR FILING DATE: 2003-02-18
24 <160> NUMBER OF SEQ ID NOS: 80
26 <170> SOFTWARE: PatentIn version 3.3
28 <210> SEQ ID NO: 1
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41 cagacccccgg agaagcgcct ggagtgggtc gccaccatca acaacggcgg ctcccacacc     180
43 tactgctccg acaacgtgaa gggccgcttc accaccttcc gcgacaacgt gaagaacacc     240
45 ctctacctcc agatgtcctc cctcaacttc gaggacaccg ccatgtacta ctgcgcccgc     300
47 gcctactacc gcttcgacgt gcgcgcctgg ttctcctact ggggccaggg caccctcgtg     360
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65 accgacggct cccccgcgcct cctcatccag tgcgcctccg agtccatctc cggcatcccg     180
67 tcccgcttct ccggcaccgg ctccggcacc gacttcaccc tcaacttcaa ctccgtggag     240
69 tccgagtaca tcaccgacta ctactgccag cagtccaaca cctggccgac ctacccgttc     300
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RAW SEQUENCE LISTING
PATENT APPLICATION: **US/10/544,284B**

DATE: 02/10/2007
TIME: 09:53:52

Input Set : **A:\corrected 70235USPCT.ST25.txt**
Output Set: **N:\CRF4\02102007\J544284B.raw**

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 73 ccgccgtcct ccgagcagct cacctccggc ggcgcgtccg tggtgtgctt cctcaacaac     420
 75 ttctacccga aggacatcaa cgtgaagtgg aagatcgacg gctccgagcg ccagaacggc     480
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 92 cagactccgg aaaagaggct ggagtgggtc gcaaccatta ataatggtgg tagtcacacc     180
 94 tactgttcag acaatgtgaa gggacgattt acaactttca gagacaatgt caaaaacacc     240
 96 ctgtaccttc aaatgagcag tctgaacttt gaggacacag ccatgtatta ctgtgcaaga     300
 98 gcctactata ggttcgacgt gagggcctgg ttttcttatt ggggccaagg gactctggtc     360
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115 ccttccaggt ttagtggcac tggatcaggg acagatttta ctcttaactt caacagtgtg     240
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137 tactacgccg acaccgtgaa gggccgcttc accgtgtccc gcgacaaccc gaagtccacc     240
139 ctcttcctcc agatgaacct cctccgcagc gaggacaccg ccatgtacta ctgcgcccgc     300
141 gacgactacg gctcctccgg ctggtacttc gacgtctggg gcgctggcac cacggtgacc     360
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159 ctcgcctggt atcagcagaa gccgggccag tccccgaagc tcctcatcta ctgggcctcc     180
161 acccgcgagt ccggcgtgcc ggaccgcttc accggctccg gctccggcac cgacttcacc     240
163 ctcaccatct cctccgtgca ggcggaggac ctcgccgtgt actactgcac ccagtcctac     300
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181 cagagccatg gagagagcct tgagtggatt ggatatatta atttttacaa tggtgctact     180
183 aactacaacc agaagttcaa gggcaaggcc acatttactg tagacacatc ctccagcaca     240
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192 <210> SEQ ID NO: 8
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205 cagaagtcag gtgcctcccc caaactctgg atttatagca catccaactt ggcttctgga     180
207 gtccctgctc gcttcagtgg cagtgggtct gggacctctt actctctcac aatcagcagt     240
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231 agtagtgtct acctgcaaat gaacaactta agagctgaag acactggcat ttattactgt     300
233 accaggcctt actacggtgg taggttcttc tactggtact tcgatgtctg gggcgcaggg     360
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251 cagaagccag gattctcccc taaactcttg atttatagga catccaatct ggcttctgga    180
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316 ttactacgga actatgctat ggactactgg ggtcaaggaa cctcagtcac cgtctcctca   360
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322 <211> LENGTH: 317
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362 <211> LENGTH: 318
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364 <213> ORGANISM: mouse
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371 ccaggattct cccctaaact cttgatttat aggacatcca atctggcttc tggagtccca     180
373 gttcgcttca gtggcagtgg gtctgggacc tcttactctc tcacaattgg caccatggag     240
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383 <213> ORGANISM: artificial sequence
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386 <223> OTHER INFORMATION: anti-K99 heavy chain variable region
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394 Pro Gly Gly Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
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398 Ser Asp Tyr Phe Met Ser Trp Ile Arg Gln Thr Pro Glu Lys Arg Leu
399         35                  40                  45
402 Glu Trp Val Ala Thr Ile Asn Asn Gly Gly Ser His Thr Tyr Cys Ser
403     50                  55                  60
406 Asp Asn Val Lys Gly Arg Phe Thr Thr Phe Arg Asp Asn Val Lys Asn
407 65                  70                  75                  80
410 Thr Leu Tyr Leu Gln Met Ser Ser Leu Asn Phe Glu Asp Thr Ala Met
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FYI

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 02/10/2007
PATENT APPLICATION: US/10/544,284B   TIME: 09:53:53

Input Set : **A:\corrected 70235USPCT.ST25.txt**
Output Set: **N:\CRF4\02102007\J544284B.raw**

<u>Please Note:</u>
**Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.**

Seq#:18; Xaa Pos. 225,226
Seq#:80; Xaa Pos. 2..

VERIFICATION SUMMARY
PATENT APPLICATION: **US/10/544,284B**

DATE: 02/10/2007
TIME: 09:53:53

Input Set : **A:\corrected 70235USPCT.ST25.txt**
Output Set: **N:\CRF4\02102007\J544284B.raw**

L:498 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:224
L:1890 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80 after pos.:0